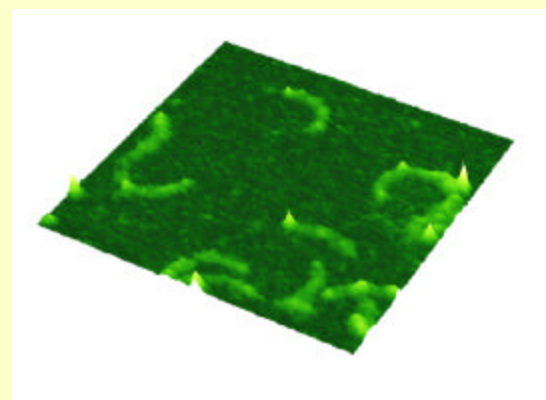
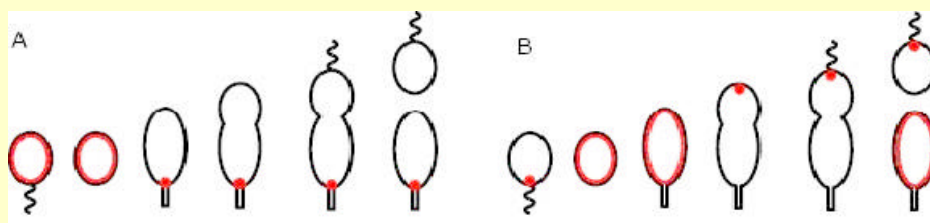


Advanced Microscopy for Protein Localization in Caulobacter



W. E. Moerner, Matt Paige, S. Nishimura
Ellen Judd, Lucy Shapiro
Stanford University
supported in part by DARPA

The Problem

- Differentiation and asymmetric cell division in *Caulobacter* have been shown to be accompanied by a pattern of protein localization

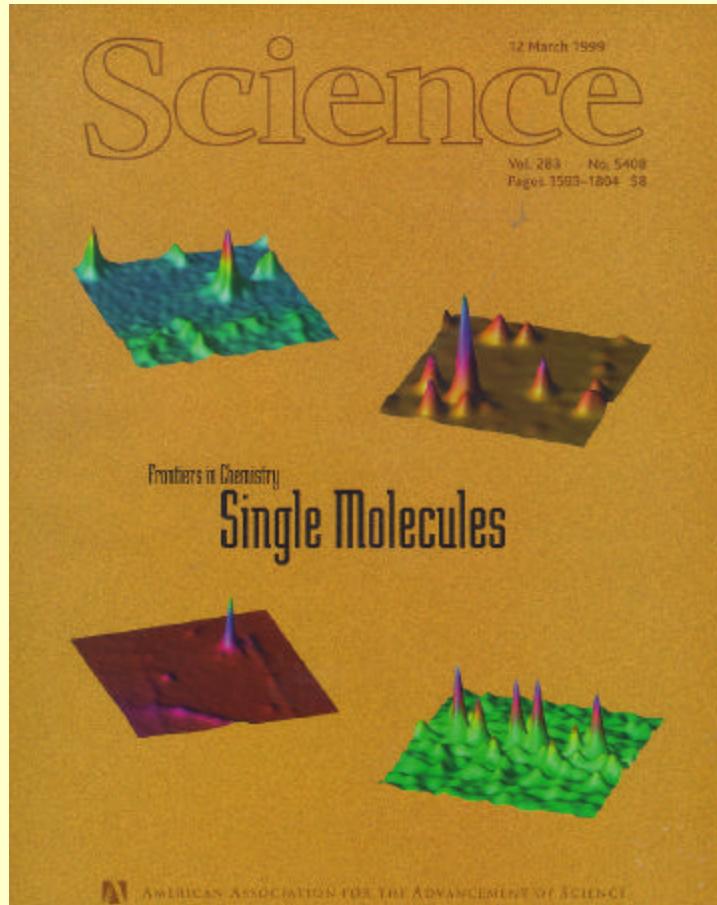
- Detecting location of proteins in live bacteria can be done by fluorescent labeling with GFP

- Research Challenge:

Determine real-time location of proteins such as DivJ, PleC, in a $1\mu\text{m} \times 2\mu\text{m}$ sized object. Requires subwavelength resolution and extreme sensitivity

- Plan: Utilize ultimate sensitivity of single-molecule microscopy coupled with selected near-field and ultraresolution microscopies

The Promise



This cover of Science (March 1999) shows imaging of individual molecules by the Moerner Lab by a variety of techniques, both at low temperatures, as well as in room temperature biomolecular environments.

Key idea: Observing individuals removes ensemble averaging, allowing exploration of

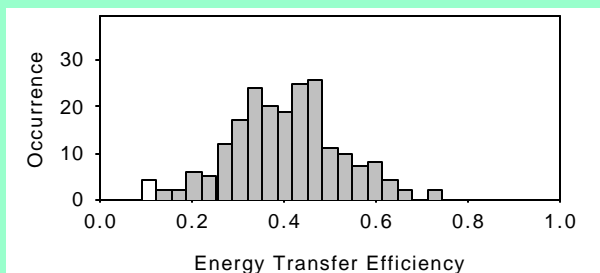
- static heterogeneity (different folds, structures, local environments, ...)
- dynamic heterogeneity (different nucleotide states, activities, ...)

Optical Study of Single Molecules (1.66 μm)

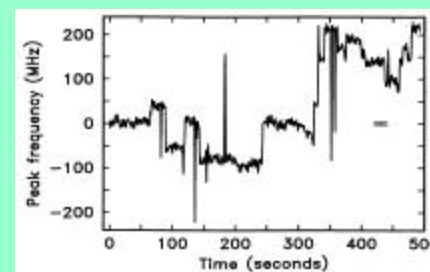
We *optically* study the spectroscopy, orientation, motion, and dynamical behavior of *single, individual* chromophores in *complex* environments: solids, liquids, proteins, ...

Why is this possibly useful?

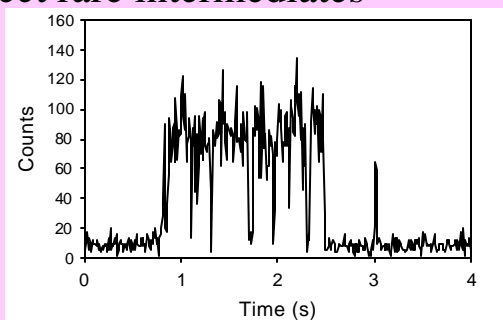
No ensemble averaging: direct observation of (static and dynamic) **heterogeneity** by measuring the full distribution



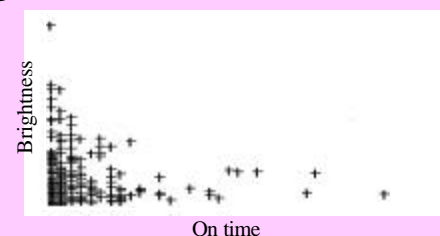
Extreme sensitivity to immediate local **nanoenvironment**; sense local strain field changes (spectral diffusion) or local optical field



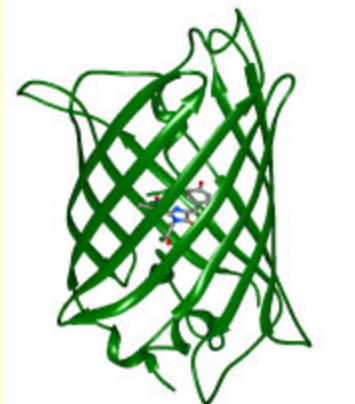
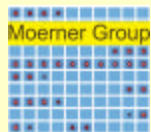
Time-dependent state changes directly observable, without synchronization, and can detect rare intermediates



Statistical **correlations** possible: can plot lifetime vs linewidth, brightness vs on-time, ...



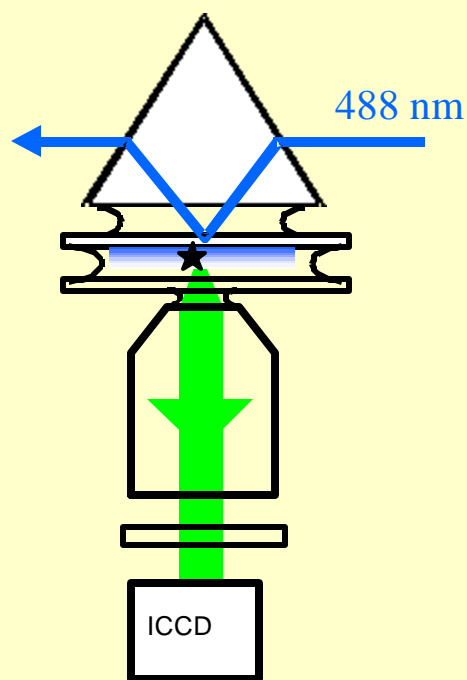
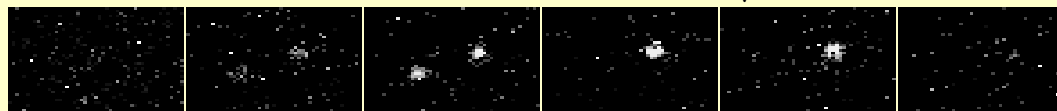
Quantum optics: photon antibunching, nonclassical light, ... for single molecule trapped by the solid



Blinking and Switching Dynamics for Single Copies of Green Fluorescent Protein



3x2 μm , 100 ms frames



Wide-field TIR microscopy,
water-filled agarose gels

Emission characteristics:
Short on-time, long off time,
reversible

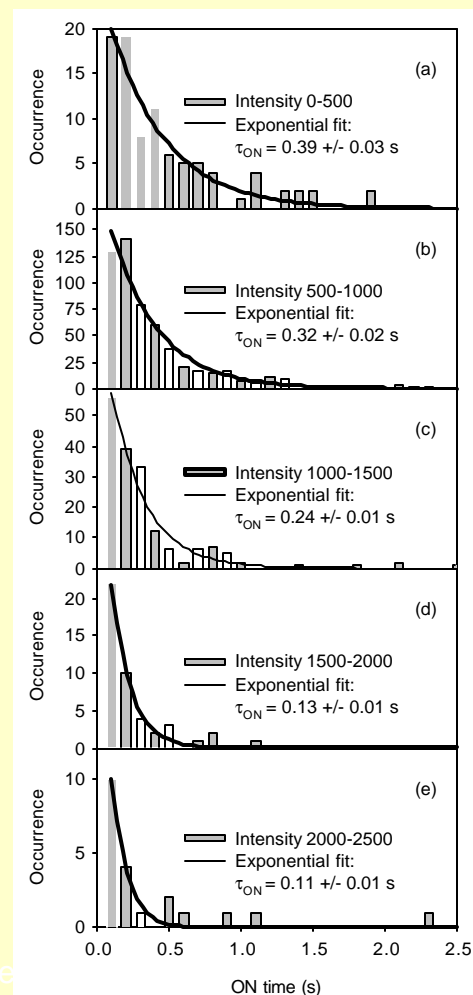
On-time distribution scales
inversely with intensity

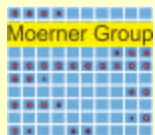
No pH dependence

Consistent with reversible
bleaching with q.e. 10^{-5}

Probable mechanism:
photoisomerization

Nature (1997) 388, 355
J. Phys. Chem. (1999) A103, 1553

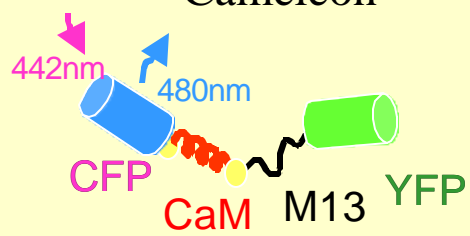




Detection of Local Calcium Concentrations by Fluorescence Resonance Energy Transfer From Single Molecules

Sophie Brasselet, Erwin Peterman, Atsushi Miyawaki*, Roger Tsien*, and W.E. Moerner

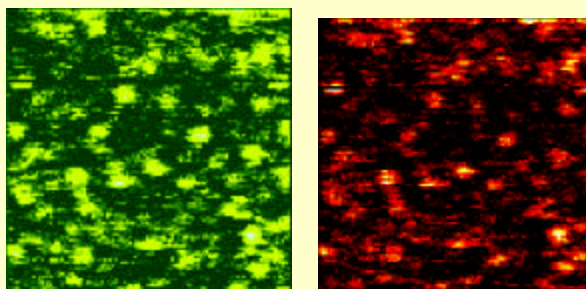
Dual-GFP
“Cameleon”



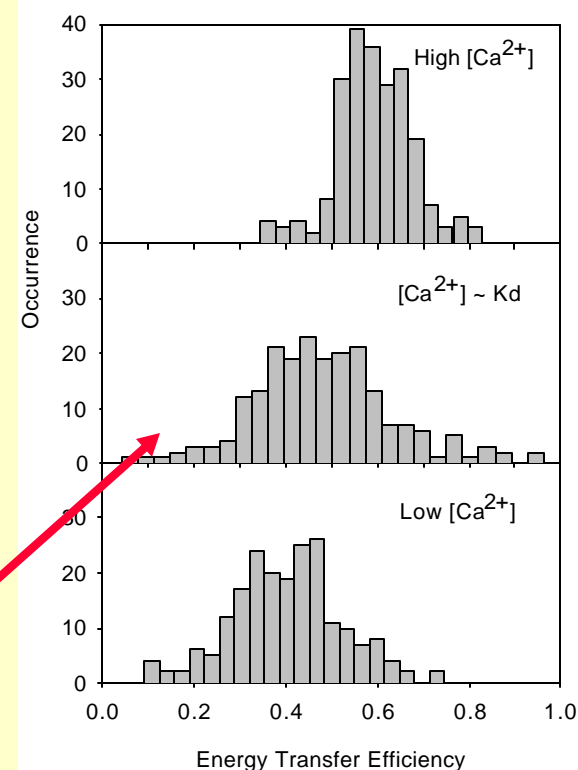
+4 Ca^{2+}



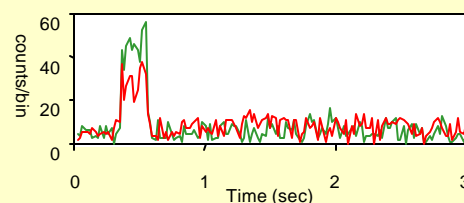
2-color confocal microscopy



single-molecule histograms



SM time traces: D and A



excess width at intermediate
[Ca⁺⁺] due to binding kinetics

*Collaborators from UC San Diego

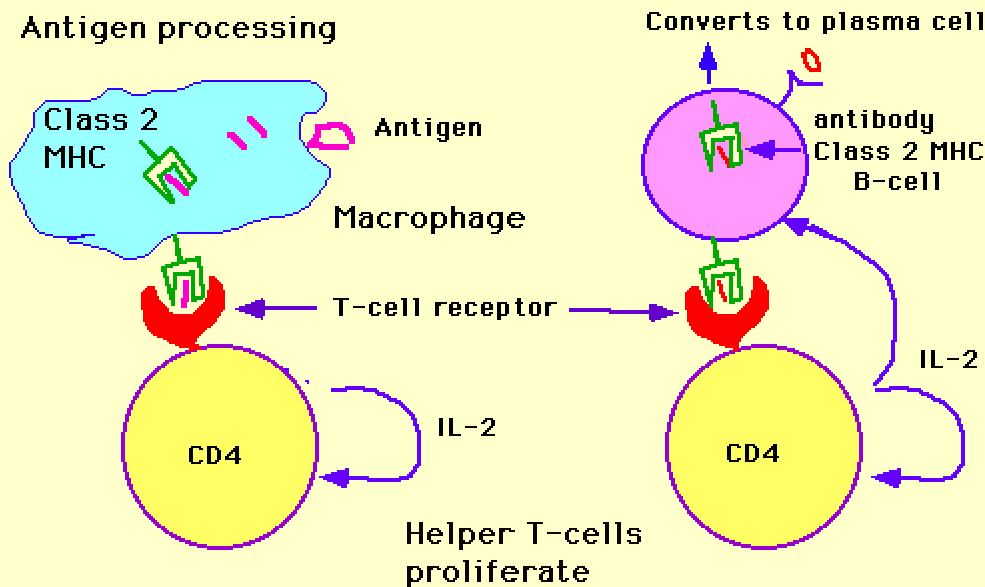
See J. Phys. Chem B 104, 3676-3682 (2000).

Imaging Single Molecules *in vivo*

Recent advance (unpublished):
Imaging of single MHCII proteins on live cells

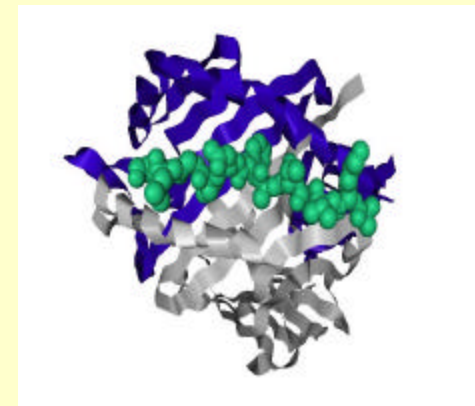
S. Brasselet, S. Nishimura, WEM
M. Vrljic, H. McConnell

Background:



from The Biology Project, developed at
The University of Arizona.

extracellular view



Fremont, et al. Science 1996, 272: 1001

Key idea: peptide off-times
are ~200 hr; label peptides
with fluorescent tag via -SH